A method of identifying variations in the starch synthesis characteristics of a cereal plant, comprising the step of

- (a) identifying a variation in nucleic acid sequence in the intron regions of the SBE I, SBE II, SSS I or DBE genes, or
- (b) identifying a variation in nucleic acid sequence compared to the sequence shown in one or more of SEQ ID NO:5, SEQ ID NO:7, SEQ ID NO:9, SEQ ID NO:10, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:15, SEQ ID NO:16, or SEQ ID NO:17.

出う プ5. A product comprising plant material propagated from a plant according to claim 78. 43

A product comprising plant material propagated from a plant in which a gene was targeted to the endosperm of a cereal plant, by a nucleic acid construct according to claim 70.

A product according to claim 76, wherein the product is a food product.

## **REMARKS**

Entry of the foregoing amendments prior to examination is respectfully requested.

Respectfully submitted,

Date March 10, 2000

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